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(4-Bromophenyl)(1*H*-indol-7-yl)-methanoneGrzegorz Dutkiewicz,^a C. S. Chidan Kumar,^b
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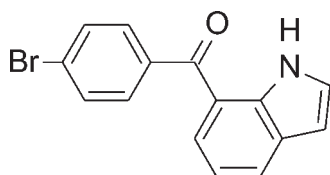
Received 1 October 2009; accepted 7 October 2009

Key indicators: single-crystal X-ray study; *T* = 291 K; mean $\sigma(\text{C}—\text{C}) = 0.003 \text{ \AA}$; *R* factor = 0.024; *wR* factor = 0.063; data-to-parameter ratio = 14.7.

In the crystal, the molecules of the title compound, $\text{C}_{15}\text{H}_{10}\text{BrNO}$, are connected into centrosymmetric dimers by pairs of $\text{N}—\text{H}\cdots\text{O}$ hydrogen bonds. The dihedral angle between the planes of indole ring system and benzene ring is $50.13(5)^\circ$. The indole plane is significantly less twisted from the plane of the central $\text{C}—\text{C}(=\text{O})—\text{C}$ bridge than the benzene plane [dihedral angles = $15.51(3)$ and $40.13(7)^\circ$, respectively]. The bond angles within the benzene ring show an approximately additive effect of the influence of both substituents.

Related literature

For applications of indoles, see: Murphy *et al.* (1997); Gupta *et al.* (1982); Al-Soud *et al.* (2004); Shigenaga *et al.* (1993); Butera *et al.* (2001). For synthetic procedures, see: Robinson (1982); Walsh *et al.* (1984). For related crystal structures of 7-pyridylindoles, see: Mudadu *et al.* (2006). For the influence of the substituent on the geometry of the phenyl ring, see: Domenicano (1988). For a description of the Cambridge Structural Database, see: Allen (2002).



Experimental

Crystal data

 $\text{C}_{15}\text{H}_{10}\text{BrNO}$
 $M_r = 300.15$
Monoclinic, $P2_1/n$
 $a = 11.3241(4) \text{ \AA}$
 $b = 7.4651(3) \text{ \AA}$ $c = 14.9579(5) \text{ \AA}$
 $\beta = 103.100(4)^\circ$
 $V = 1231.57(8) \text{ \AA}^3$
 $Z = 4$
Mo $K\alpha$ radiation $\mu = 3.32 \text{ mm}^{-1}$
 $T = 291 \text{ K}$ $0.4 \times 0.2 \times 0.15 \text{ mm}$

Data collection

Oxford Diffraction Xcalibur
(Sapphire2, large Be window)
diffractometer
Absorption correction: multi-scan
(*CrysAlis Pro*; Oxford)Diffraction, 2009)
 $T_{\min} = 0.26$, $T_{\max} = 0.60$
25357 measured reflections
2558 independent reflections
1864 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.036$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.024$
 $wR(F^2) = 0.063$
 $S = 1.06$
2558 reflections174 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.34 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.29 \text{ e \AA}^{-3}$

Table 1

Hydrogen-bond geometry (\AA , $^\circ$).

$D—H\cdots A$	$D—H$	$H\cdots A$	$D\cdots A$	$D—H\cdots A$
$\text{N1}—\text{H1}\cdots\text{O10}^i$	0.86	2.14	2.935 (2)	153

Symmetry code: (i) $-x + 1, -y, -z + 2$.

Data collection: *CrysAlis Pro* (Oxford Diffraction, 2009); cell refinement: *CrysAlis Pro*; data reduction: *CrysAlis Pro*; program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1993); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *Stereochemical Workstation Operation Manual* (Siemens, 1989); software used to prepare material for publication: *SHELXL97*.

CSC thanks the University of Mysore for research facilities.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: ER2074).

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supporting information

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(4-Bromophenyl)(1*H*-indol-7-yl)methanone

Grzegorz Dutkiewicz, C. S. Chidan Kumar, H. S. Yathirajan and Maciej Kubicki

S1. Comment

The synthesis of indole derivatives has long been a topic of fundamental interest to organic and medicinal chemists (Murphy *et al.*, 1997). Indole derivatives are important pharmacologically, possessing anti-allergic (Shigenaga *et al.*, 1993), central-nervous-system depressant (Sen Gupta *et al.*, 1982), muscle relaxant (Butera *et al.*, 2001), and anti-cancer (Al-Soud *et al.*, 2004) properties. The Fischer indole synthesis is the most widely used method for the preparation of indole derivatives (*e.g.*, Robinson, 1982). The title compound (I) is an intermediate for preparation of bromofenac, which is used as analgesic.

There are only few crystal structures of 7-substituted indoles in the Cambridge Crystallographic Database (Allen, 2002). Recently, the crystal structures of three 7-pyridylindoles (Mudadu *et al.*, 2006) have been reported.

The conformation of the molecule I can be described by the mutual orientation of the three approximately planar fragments (Fig. 1): indole system (maximum deviation from the least-squares plane of 0.0142 (7) Å), phenyl ring (maximum deviation 0.0145 (13) Å), and the central C—C(=O)—C bridge (0.0040 (16) Å). The dihedral angle between the terminal planes, of indole and phenyl fragments, is 50.13 (5)°, while it can be noted that the indole plane is less inclined with respect to the central bridge plane (15.51 (3)°) than is the phenyl one (40.13 (7)°). The geometry of the phenyl ring is affected by the presence of substituents; using the values given by Domenicano (1988) and obtained from the search of the CSD (Allen, 2002), it might be shown that the overall influence on the bond angles pattern is close to additivity of separate effects of both Br and COAr substituents.

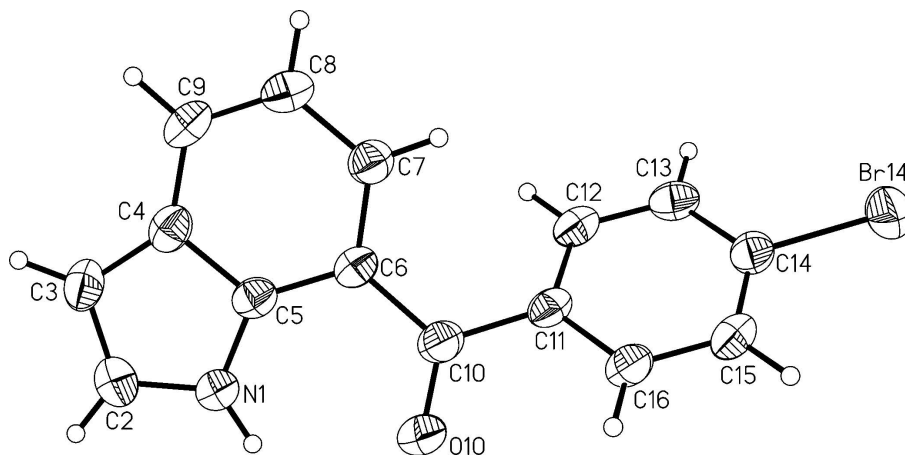
In the crystal structure the molecules of (I) are connected into the centrosymmetric, hydrogen bonded pairs - $R^2_2(12)$ motifs - by means of relatively strong and linear N—H \cdots O hydrogen bonds (Fig. 2). These dimers are packed by means of van der Waals and weak C—H \cdots π interactions.

S2. Experimental

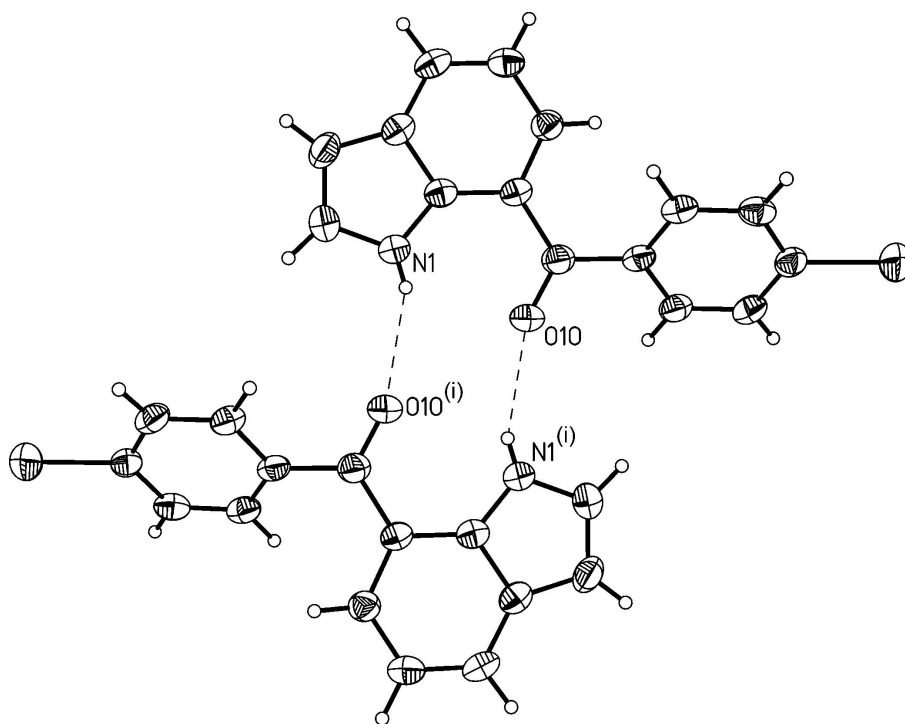
A mixture of (4-bromophenyl)(2,3-dihydro-1*H*-indol-7-yl)methanone (2.4 g, 7.9 mmol) and activated manganese dioxide (2.2 g, 25 mmol) in 100 ml dichloromethane was refluxed for 18 h (Fig. 3). The contents were filtered and the organic layer was concentrated. The product formed (Walsh *et al.*, 1984) was crystallized from tetrahydrofuran (m.p.: 435 – 437 K).

S3. Refinement

Hydrogen atoms were put in the idealized positions, and refined as riding model.

**Figure 1**

Anisotropic ellipsoid representation of the compound **I** together with atom labelling scheme. The ellipsoids are drawn at 50% probability level, hydrogen atoms are depicted as spheres with arbitrary radii.

**Figure 2**

The hydrogen bonded dimer of molecules of **I**. Hydrogen bonds are shown as dashed lines. Symmetry code: (i) 1 - x, -y, 2 - z.

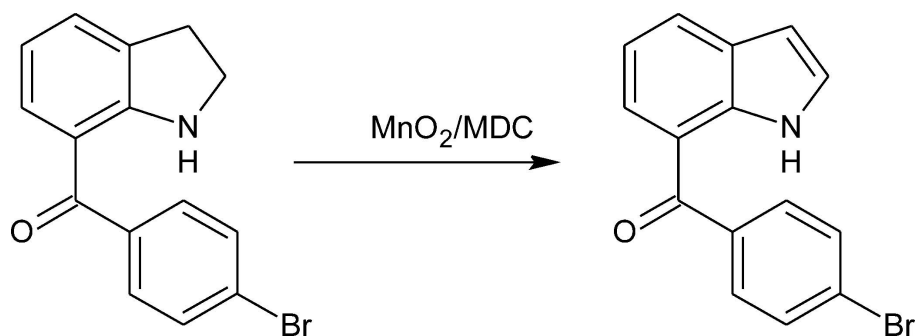


Figure 3

The preparation of the title compound.

(4-Bromophenyl)(1H-indol-7-yl)methanone

Crystal data

$\text{C}_{15}\text{H}_{10}\text{BrNO}$

$M_r = 300.15$

Monoclinic, $P2_1/n$

Hall symbol: -P 2yn

$a = 11.3241$ (4) Å

$b = 7.4651$ (3) Å

$c = 14.9579$ (5) Å

$\beta = 103.100$ (4)°

$V = 1231.57$ (8) Å³

$Z = 4$

$F(000) = 600$

$D_x = 1.619$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 12183 reflections

$\theta = 2.0\text{--}26.8^\circ$

$\mu = 3.32$ mm⁻¹

$T = 291$ K

Block, colourless

$0.4 \times 0.2 \times 0.15$ mm

Data collection

Oxford Diffraction Xcalibur (Sapphire2, large Be window)

diffractometer

Radiation source: Enhance (Mo) X-ray Source

Graphite monochromator

Detector resolution: 8.1929 pixels mm⁻¹

ω scans

Absorption correction: multi-scan

(*CrysAlis PRO*; Oxford Diffraction, 2009)

$T_{\min} = 0.26$, $T_{\max} = 0.60$

25357 measured reflections

2558 independent reflections

1864 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.036$

$\theta_{\max} = 26.8^\circ$, $\theta_{\min} = 2.1^\circ$

$h = -13 \rightarrow 14$

$k = -9 \rightarrow 9$

$l = -18 \rightarrow 18$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.024$

$wR(F^2) = 0.063$

$S = 1.06$

2558 reflections

174 parameters

0 restraints

Primary atom site location: structure-invariant direct methods

Secondary atom site location: difference Fourier map

Hydrogen site location: inferred from neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.035P)^2]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.001$

$\Delta\rho_{\max} = 0.34$ e Å⁻³

$\Delta\rho_{\min} = -0.29$ e Å⁻³

Extinction correction: *SHELXL97* (Sheldrick, 2008), $F_c^* = kF_c[1 + 0.001x F_c^2 \lambda^3 / \sin(2\theta)]^{-1/4}$

Extinction coefficient: 0.0139 (9)

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
N1	0.28642 (14)	−0.0239 (2)	0.94540 (10)	0.0386 (4)
H1	0.3601	−0.0274	0.9768	0.053 (6)*
C2	0.18656 (19)	−0.0579 (3)	0.97896 (14)	0.0459 (5)
H2	0.1876	−0.0872	1.0396	0.043 (5)*
C3	0.08506 (18)	−0.0424 (3)	0.91063 (14)	0.0449 (5)
H3	0.0056	−0.0604	0.9157	0.053 (6)*
C4	0.12356 (15)	0.0064 (3)	0.83002 (13)	0.0344 (4)
C5	0.25153 (15)	0.0167 (3)	0.85411 (11)	0.0315 (4)
C6	0.32088 (15)	0.0562 (2)	0.78996 (12)	0.0314 (4)
C7	0.25687 (16)	0.0912 (2)	0.70024 (12)	0.0353 (4)
H7	0.2995	0.1200	0.6559	0.035 (5)*
C8	0.13073 (17)	0.0839 (2)	0.67577 (13)	0.0400 (5)
H8	0.0908	0.1083	0.6155	0.046 (6)*
C9	0.06422 (17)	0.0413 (2)	0.73904 (14)	0.0387 (5)
H9	−0.0199	0.0357	0.7215	0.053 (6)*
C10	0.45391 (16)	0.0642 (2)	0.81808 (12)	0.0372 (4)
O10	0.50404 (12)	0.0723 (2)	0.90015 (9)	0.0593 (5)
C11	0.53044 (15)	0.0650 (2)	0.74900 (12)	0.0322 (4)
C12	0.50633 (16)	−0.0445 (2)	0.67135 (13)	0.0352 (4)
H12	0.4387	−0.1188	0.6605	0.041 (6)*
C13	0.58143 (17)	−0.0438 (2)	0.61058 (12)	0.0371 (5)
H13	0.5656	−0.1189	0.5596	0.035 (5)*
C14	0.68042 (16)	0.0694 (3)	0.62617 (12)	0.0363 (4)
Br14	0.779765 (18)	0.07710 (3)	0.539996 (14)	0.05633 (12)
C15	0.70737 (16)	0.1777 (3)	0.70285 (12)	0.0389 (5)
H15	0.7746	0.2529	0.7129	0.051 (6)*
C16	0.63296 (15)	0.1728 (3)	0.76455 (12)	0.0366 (4)
H16	0.6519	0.2428	0.8173	0.036 (5)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N1	0.0302 (9)	0.0511 (11)	0.0332 (8)	−0.0010 (8)	0.0045 (7)	0.0004 (7)
C2	0.0452 (12)	0.0568 (14)	0.0390 (10)	−0.0020 (10)	0.0167 (9)	−0.0002 (10)
C3	0.0309 (11)	0.0563 (14)	0.0504 (12)	−0.0031 (10)	0.0157 (9)	−0.0039 (10)
C4	0.0294 (10)	0.0291 (10)	0.0446 (10)	0.0011 (8)	0.0080 (8)	−0.0049 (9)

C5	0.0303 (10)	0.0275 (10)	0.0350 (10)	0.0002 (8)	0.0034 (7)	−0.0032 (8)
C6	0.0268 (9)	0.0303 (11)	0.0353 (9)	−0.0016 (8)	0.0032 (7)	−0.0017 (8)
C7	0.0326 (10)	0.0355 (11)	0.0367 (10)	−0.0008 (9)	0.0057 (8)	0.0033 (8)
C8	0.0348 (11)	0.0415 (12)	0.0387 (10)	0.0027 (9)	−0.0020 (8)	0.0028 (9)
C9	0.0248 (10)	0.0376 (12)	0.0502 (11)	0.0015 (8)	0.0011 (8)	−0.0024 (9)
C10	0.0320 (10)	0.0415 (12)	0.0358 (10)	−0.0035 (9)	0.0028 (8)	0.0037 (9)
O10	0.0339 (8)	0.1051 (14)	0.0347 (7)	−0.0108 (8)	−0.0010 (6)	0.0059 (8)
C11	0.0240 (9)	0.0341 (10)	0.0353 (9)	0.0015 (8)	−0.0004 (7)	0.0034 (8)
C12	0.0283 (10)	0.0321 (12)	0.0410 (10)	−0.0031 (8)	−0.0007 (8)	0.0021 (8)
C13	0.0354 (11)	0.0350 (12)	0.0363 (10)	0.0059 (9)	−0.0015 (8)	−0.0007 (9)
C14	0.0283 (10)	0.0428 (12)	0.0366 (9)	0.0071 (9)	0.0047 (8)	0.0086 (9)
Br14	0.04658 (16)	0.0801 (2)	0.04560 (15)	0.00255 (12)	0.01738 (10)	0.00603 (11)
C15	0.0268 (10)	0.0424 (12)	0.0454 (11)	−0.0057 (9)	0.0034 (8)	−0.0012 (9)
C16	0.0284 (10)	0.0402 (12)	0.0386 (10)	−0.0011 (9)	0.0020 (8)	−0.0045 (9)

Geometric parameters (Å, °)

N1—C2	1.361 (3)	C8—H8	0.9300
N1—C5	1.367 (2)	C9—H9	0.9300
N1—H1	0.8600	C10—O10	1.232 (2)
C2—C3	1.359 (3)	C10—C11	1.492 (2)
C2—H2	0.9300	C11—C16	1.388 (2)
C3—C4	1.419 (3)	C11—C12	1.395 (3)
C3—H3	0.9300	C12—C13	1.378 (3)
C4—C9	1.399 (3)	C12—H12	0.9300
C4—C5	1.414 (2)	C13—C14	1.381 (3)
C5—C6	1.402 (2)	C13—H13	0.9300
C6—C7	1.398 (2)	C14—C15	1.380 (3)
C6—C10	1.471 (2)	C14—Br14	1.8941 (18)
C7—C8	1.393 (2)	C15—C16	1.384 (2)
C7—H7	0.9300	C15—H15	0.9300
C8—C9	1.374 (3)	C16—H16	0.9300
C2—N1—C5	109.49 (16)	C8—C9—C4	119.72 (17)
C2—N1—H1	125.3	C8—C9—H9	120.1
C5—N1—H1	125.3	C4—C9—H9	120.1
C3—C2—N1	109.79 (18)	O10—C10—C6	119.87 (17)
C3—C2—H2	125.1	O10—C10—C11	118.77 (17)
N1—C2—H2	125.1	C6—C10—C11	121.36 (15)
C2—C3—C4	106.88 (17)	C16—C11—C12	118.51 (17)
C2—C3—H3	126.6	C16—C11—C10	118.77 (16)
C4—C3—H3	126.6	C12—C11—C10	122.66 (16)
C9—C4—C5	118.45 (17)	C13—C12—C11	120.89 (17)
C9—C4—C3	134.58 (18)	C13—C12—H12	119.6
C5—C4—C3	106.97 (16)	C11—C12—H12	119.6
N1—C5—C6	130.57 (15)	C12—C13—C14	119.29 (17)
N1—C5—C4	106.86 (16)	C12—C13—H13	120.4
C6—C5—C4	122.53 (15)	C14—C13—H13	120.4

C7—C6—C5	116.57 (16)	C15—C14—C13	121.14 (17)
C7—C6—C10	122.80 (16)	C15—C14—Br14	119.62 (14)
C5—C6—C10	120.61 (15)	C13—C14—Br14	119.24 (14)
C8—C7—C6	121.52 (17)	C14—C15—C16	119.04 (18)
C8—C7—H7	119.2	C14—C15—H15	120.5
C6—C7—H7	119.2	C16—C15—H15	120.5
C9—C8—C7	121.18 (17)	C15—C16—C11	121.07 (18)
C9—C8—H8	119.4	C15—C16—H16	119.5
C7—C8—H8	119.4	C11—C16—H16	119.5
C5—N1—C2—C3	0.7 (2)	C3—C4—C9—C8	−179.0 (2)
N1—C2—C3—C4	−0.8 (2)	C7—C6—C10—O10	163.69 (18)
C2—C3—C4—C9	179.9 (2)	C5—C6—C10—O10	−14.7 (3)
C2—C3—C4—C5	0.6 (2)	C7—C6—C10—C11	−15.5 (3)
C2—N1—C5—C6	−178.08 (19)	C5—C6—C10—C11	166.12 (17)
C2—N1—C5—C4	−0.3 (2)	O10—C10—C11—C16	−37.9 (3)
C9—C4—C5—N1	−179.61 (17)	C6—C10—C11—C16	141.34 (18)
C3—C4—C5—N1	−0.2 (2)	O10—C10—C11—C12	139.08 (19)
C9—C4—C5—C6	−1.6 (3)	C6—C10—C11—C12	−41.7 (3)
C3—C4—C5—C6	177.79 (17)	C16—C11—C12—C13	−1.0 (3)
N1—C5—C6—C7	179.52 (18)	C10—C11—C12—C13	−177.97 (16)
C4—C5—C6—C7	2.0 (3)	C11—C12—C13—C14	−1.2 (3)
N1—C5—C6—C10	−2.0 (3)	C12—C13—C14—C15	2.0 (3)
C4—C5—C6—C10	−179.52 (18)	C12—C13—C14—Br14	−177.47 (13)
C5—C6—C7—C8	−1.1 (3)	C13—C14—C15—C16	−0.4 (3)
C10—C6—C7—C8	−179.52 (17)	Br14—C14—C15—C16	179.00 (14)
C6—C7—C8—C9	−0.2 (3)	C14—C15—C16—C11	−1.9 (3)
C7—C8—C9—C4	0.7 (3)	C12—C11—C16—C15	2.5 (3)
C5—C4—C9—C8	0.2 (3)	C10—C11—C16—C15	179.66 (17)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
N1—H1 \cdots O10 ⁱ	0.86	2.14	2.935 (2)	153

Symmetry code: (i) $-x+1, -y, -z+2$.